	-	N						4		9	
	7	2						6		0	
HDVAM	GREDILE	GREDILEQWVSGRKKLEELERDLRKLKKKIKKLEEDNP <b>WL</b> GNIKGIIGK	KLEE	LERDL	RKLKI	KKIK	KLEE]	DNPWI	GNIK	BIIGK	
HDVL1	GREEVLE	GREEVLEQWVNSRKKAEELERDLRKTKKKIKKLEDDNP <b>WL</b> GNIKGILGK	KAEE	LERDL	RKTKI	KKIK	KLED.	DNPWI	GNIKC	BILGK	
HDVD3	GREEVLE	GREEVLEQWVSGRKKLEELERDLRKVKKKIKKLEDEHP <b>WL</b> GNIKGILGK	KLEEI	LERDL	RKVKI	KKIK	KLED	EHPWI	GNIKC	SILGK	
HDVNA	GREEVLE	GREEVLEQWVAGRRKQEELERDLRKTKKKIKKLEEENP <b>WL</b> GNIKGILGK	KQEEJ	ERDL	RKTKI	KKIK	KLEE	ENPW	GNIK	BILGK	
HDVS1	TREETLE	TREETLEKWITARKKAEELEKDLRKTRKTIKKLEEENP <b>WL</b> GNIVGII.R	KAEEI	LEKDL	RKTRI	KTIK	KLEE	ENPW	GNIVC	BII.R	
HDVS2	TREETLE	TREETLEKWITARKKAEELEKDLRKARKTIKKLEEENPWLGNILGII.R	KAEEI	LEKDL	RKARI	<b>KTIK</b>	KLEE	ENPW	GNILC	SII.R	
HDVM1	GREQILE	GREQILEQWVDGRKKLEELERDLRKIKKKIKKLEEENP <b>WL</b> GNVKGILGK	KLEE	LERDL	RKIKI	KKIK	KLEE	ENPW	GNVKC	SILGK	
HDVM2	GREQILE	GREQILEQWVDGRKKLEELERDLRKIKKKIKKLEEENP <b>WL</b> GNVKGILGK	KLEE	LERDL	RKIKI	KKIK	KLEE	ENPW	GNVKC	SILGK	
HDVIT	GREEILE	GREEILEQWVAGRKKLEELERDLRKTKKKLKKIEDENP <b>WL</b> GNIKGILGK	KLEE	LERDL	RKTKI	KKLK	KIED	ENPW	GNIK	SILGK	
HDVWO	GREEILE	GREEILEQWVAGRKKLEELERDLRKTKKKLKKIEDENP <b>WL</b> GNIKGILGK	KLEE	LERDL	RKTKI	KKLK	KIED	ENPW	GNIKC	BILGK	
	ъ *	\$\d\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\	*	*	*	*	*	ჯ	ש	Д	

Figure 1

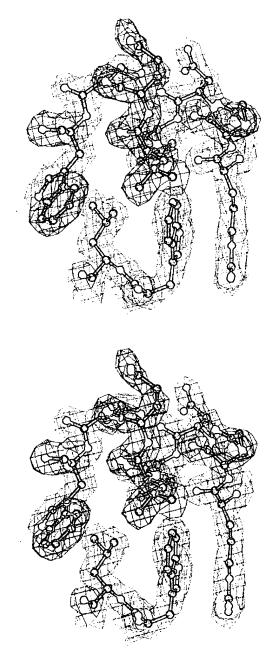
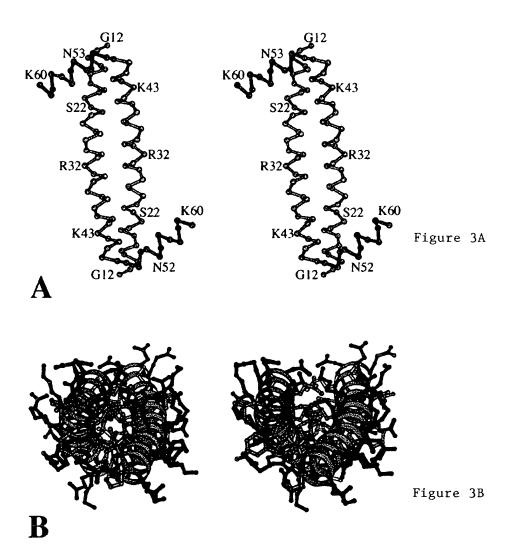


Figure 2



12 cahcdefca

48

gabcdefgabcdefgabcdefgabcdefga
GREDILEQWVSGRKKLEELERDLRKLKKKIKKLEEDN
NDEELKKIKKKLKRLDRELEELKKRGSVWQELIDERG
agfedcbagfedcbagfedcbagfedcbag
48

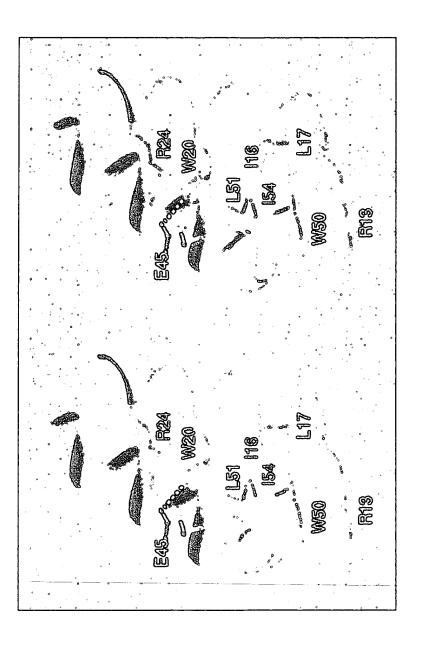


Figure 4

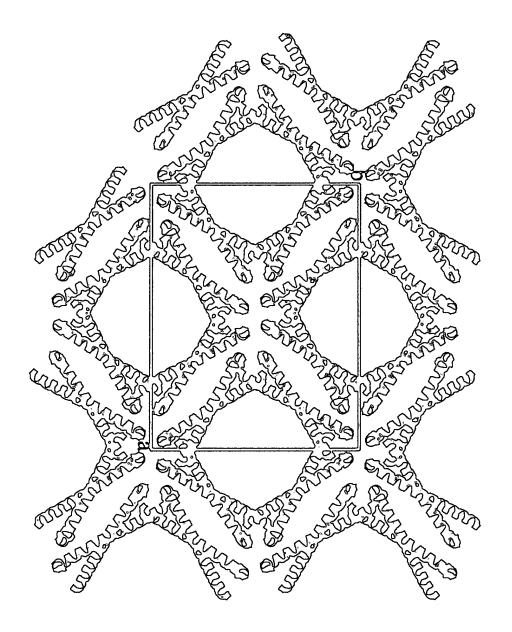


Figure 5

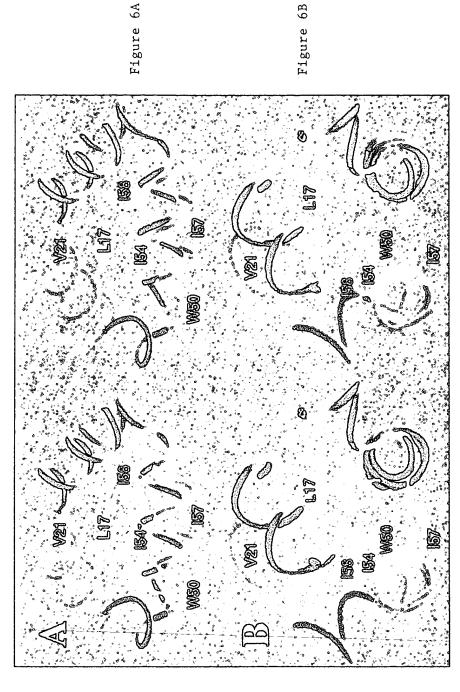


Figure 6A

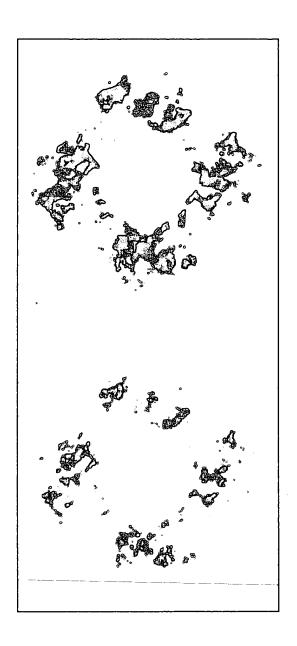
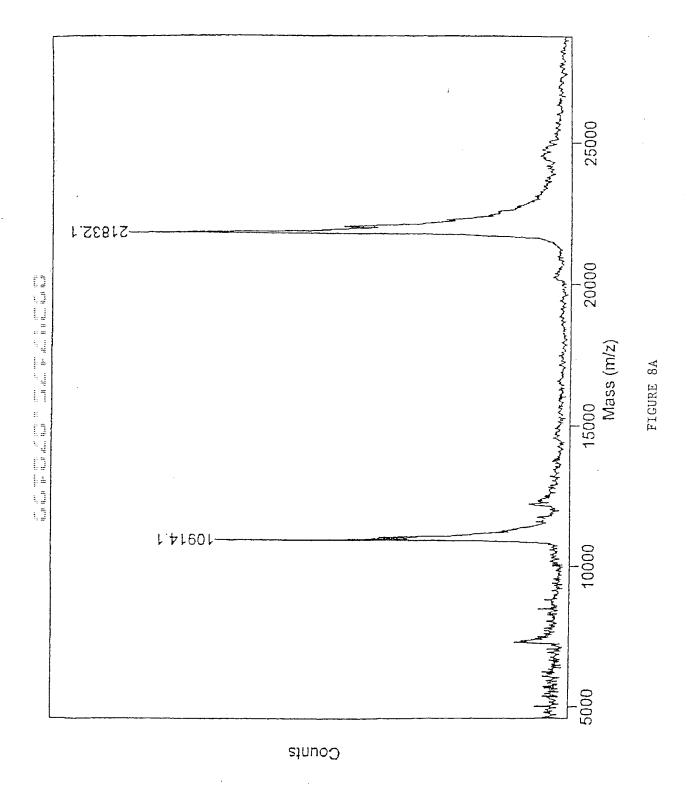


Figure 7A

Figure 7B





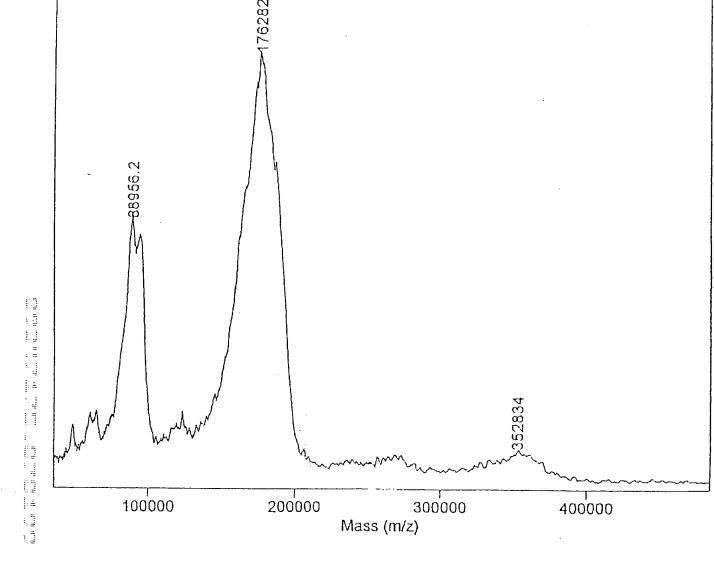


FIGURE 8B

## NdeI

- M S R S E R R K D R G  $^{\prime}$ G R E D I L E GGGCATATGAGCCGTAACGTCGTAAAGATCGTGGCGCACTTCTATAAGACCTT CTGGAACCTTCTATAAGACCTT
- Q W V S G R K K L E E L E R D L R K L K CAGTGGGTGAGCGGCCGTAAGAGTTAGAGGAATTGGAACGTGATCTGCGTAAACTGAAA GTCACCCACTCGCCGGCATTCTTCAATCTCCTTAACCTTGCACTAGACGCATTTGACTTT
- K K I K K L E E D N P W L G N I K G I I AAGAAGATTAAGAAACTGGAAGAAGATAACCCGTGGTTGGGTAATATTAAAGGCATTATT TTCTTCTATTGGGCACCAACCCATTATAATTTCCGTAATAA

- G K S L S R E E E E L K R L T E E D E  $\frac{GGCAAATCTCTGAGCCGTGAAGAAGAAGAAGAAGAAGAACTGAAACGTCTGACCGAAGAAGAAGAAGAACACTTTAGAGACTCGGCACTTCTTCTTCTTCTTGACTTTGAGACTGGCTTCTTCTACTT$
- K R E R R I A G P S V G G V N P L E G G AAACGTGAACGTCGTATTGCAGGTCCATCTGTTGGTGGTGTGAACCCGCTGGAAGGCGGC
  TTTGCACTTGCAGCATAACGTCCAGGTAGACAACCACCACACTTGGGCGACCTTCCGCCG
- S R G A P G G G F V P S M Q G V P E S P AGCCGTGGTGCACCGGGCGGTGGCTTTGTGCCGTCTATGCAAGGTGTTCCAGAAAGCCCGTCGGCACCGAAACACGGCAGATACGTTCCACAAGGTCTTTCGGGC
- F A R T G E G L D I R G S Q G F P NCOI TTTGCGCGTACCGGCGAAGGCCTGGATATTCGTGGCAGCCAGGGCTTTCCGTAAACCATGGCGC AAACGCGCATGGCCGCTTCCGGACCTATAAGCACCGTCGGTCCCGAAAGGCATTTGGTACCGCG

Figure 9

٦	G.T.AG.GGA
116	GTCP. G
233	AG
3 1 9	GIAAAAAAAAAA.
. 19 7	GGACGAGTGAGCTTATCCCCGGGAATTCCACTTATCGTCCCCCACACACGCCCCGGACCCTTCAAAGCGACCGAGGGGGGGG
581	ATGGGATECTCCTCCGAATTCCGAAATTCCTTCCCCCAAAGGTCGCCCAAGAATTGGCGGAACCCAAAGGTTCCAATCTTTCTT
669	GGTCCCAGCCTCCTCGCTGGCGCGGCGGCGAACATTCCGAAGGAACCOTCCCTCGGFAATGGCAACGCAACGCAACATCTCTTAGCTTCCTAAGATGCGAAGAAAAAG
815	TWOCTET COCTT GGOCK, TOCGAGT WORLD COTOC TOCK TOCK TOCK TOCK TOCK TOCK TOC
932	A CONTRACTEGAMACCCCCTTTATTCACTCGGGGTCGACAACAACAACAACAACACGCTGGGAAGATAAATCCCATCCAT
1049	CCTCCCCGGTTCCTTCCAAGGGGAACTCCGGGAACTCCTTGAAGGAAG
1166	CGGCTATTCPTCTCCCTCTCTCCTCGCTTCTCTCCTCCTCCCTCCC
1273	CCTTCCTTCGTGGTGATCCTCCTTCTCGTGAATCCTCCCTGAAAGGCCTTCTCTAACGTCGAATCTCTAATCTCAATCTTCTAATCTTCTAATCTTCAATCTTCGAATCTTCTCCCCGGTGATCCTCCCTGGAAAGGAATCTTCTCTAATCTAATCTTCAATCAA
1400	
7181	CCTCTALCTPOTTCPTCCCTCACCACTCCTCCCCACCTCCTCCCCCCCCTCTTTCCTCC
1634	

Figure 10

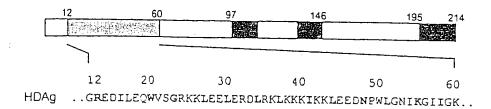
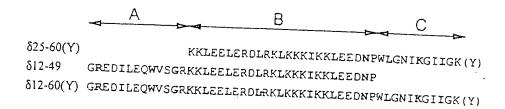


Figure 1!A



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Figure 11B

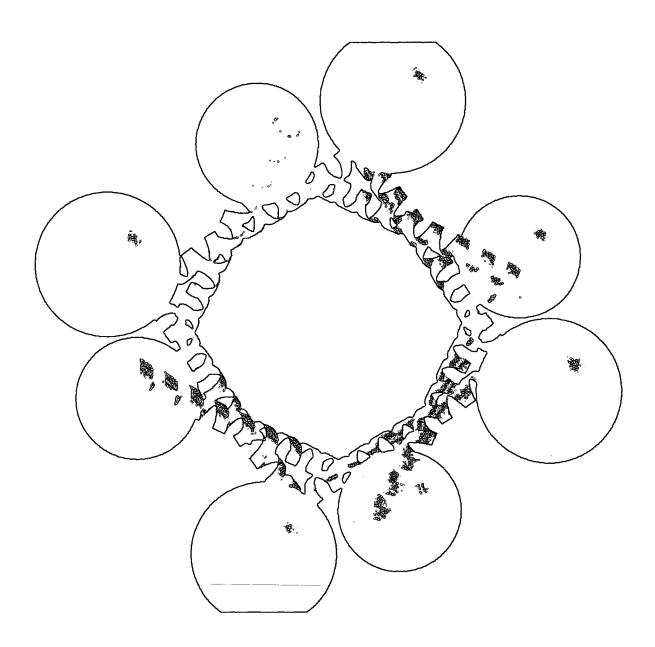
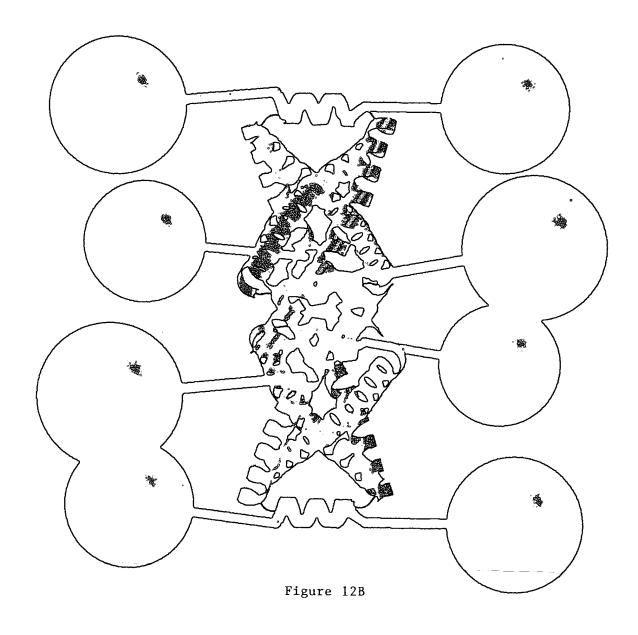
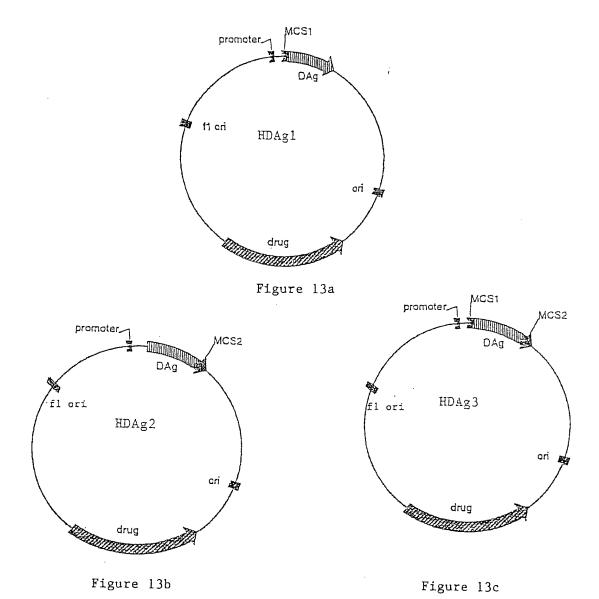


Figure 12A





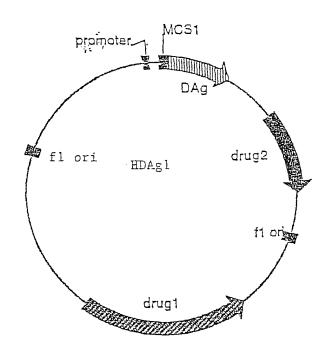


Figure 14

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1, 🗓 1, 2

Figure 15

1	GGGCATATGA	GCCGTAGCGA	ACGTCGTAAA	GATCGTGGCG	GCCGTGAAGA
51	TATTCTGGAA	CAGTGGGTGA	GCGGCCGTAA	GAAGTTAGAG	GAATTGGAAC
101	GTGATCTGCG	TAAACTGAAA	AAGAAGATTA	AGAAACTGGA	AGAAGATAA
151	CCGTGGTTGG	GTAATATTAA	AGGCATTATT	GGCAAGAAAG	ATAAAGATGO
201	CGAAGGCGCG	CCGCCGGCGA	AGAAACTGCG	TATGGATCAG	ATGGAAATTG
251	ATGCGGGCCC	GCGTAAACGT	CCGCTGCGTG	GCGGCTTTAC	CGATAAGGAA
301	CGTCAGGACC	ATCGTCGTCG	TAAAGCGCTG	GAAAACAAAC	GTAAACAGCT
351	GAGCAGCGGC	GGCAAATCTC	TGAGCCGTGA	AGAAGAAGAA	GAACTGAAAC
401	GTCTGACCGA	AGAAGATGAA	AAACGTGAAC	GTCGTATTGC	AGGTCCATCT
451	GTTGGTGGTG	TGAACCCGCT	GGAAGGCGGC	AGCCGTGGTĠ	CACCGGGCGG
501	TGGCTTTGTG	CCGTCTATGC	AAGGTGTTCC	AGAAAGCCCG	TTTGCGCGTA
551	CCGGCGAAGG	CCTGGATATT	CGTGGCAGCC	AGGGCTTTCC	GTAAACCATG
60 <u>1</u>	GCGC				•

Figure 16

	1				48
wildtype HDAg-S	MSRSERRK DI	RGGREDILE Q	WVSGRKKLE E	LERDLRKLK K	KIKKLEEDN
pR5DV5 plasmid	MSRSERRK D				
Identity	MSRSERRK D				
	49				98
wildtype HDAg-S	PWLGNIKGII	GKKDKDGEGA	PPAKKLRMDQ	MEIDAGPRKR	PLRGGFTDKE
pRSDV5 plasmid	PWLGNIKGII				PLRGGFTDKE
Identity	PWLGNIKGII				PLRGGFTDKE
	99				148
wildtype HDAg-S	RQDHRRRKAL	ENKRKOLSSG	GKSLSREEEE	ELKRLTEEDE	
pR5DV5 plasmid	RQDHRRRKAL				
Identity					KRERRIAGPS
	149				195
wildtype HDAg-S	VGGVNPLEGG	SRGAPGGGFV	PSMQGVPESP	FARTGEGLDI	RGSQGFP
pR5DV5 plasmid	VGGVNPLEGG	SRGAPGGGFV	PSMQGVPESP	FARTGEGLDI	RGSQGFP
Identity	VGGVNPLEGG	SRGAPGGGFV	PSMQGVPESP	FARTGEGLDI	RGSQGFP

Figure 17

primerl

GGGCATATGAGCCGTAGCGAACGTCGTAAAGATCGTGGCGGCCGTGAAGATA TTCTGGAACAGTGGGTGAGCGGCCGTAAGAAGTTAGAGGAA

primer2

ATATTACCCAACCACGGGTTATCTTCTTCCAGTTTCTTAATCTTCTTTTCCAGTTTACGCAGATCACGTTCCAATTCCTCTAACTTCTTACGGCC

primer3

primer4

GATGGTCCTGACGTTATCGGTAAAGCCGCCACGCAGCGGACGTTTA CGCGGGCCCGCATCAATTTCCATCTGATCCATACGCAGTTTCTT

primer5

primer6.

 ${\tt CAACAGATGGACCTGCAATACGACGTTCACGTTTTTCATCTTCTTCGGTC} \\ {\tt AGACGTTTCAGTTCTTCTTCTTCACGGCTCAGAGAT} \\$ 

primer7

TATTGCAGGTCCATCTGTTGGTGGTGTGAACCCGCTGGAAGGCGGCAGCC GTGGCGCGCGGCGGCGGCTTTGTGCCGTCTATGCAAGGTGTTCCAGAA A

primer8

GCGCCATGGTTTACGGAAAGCCCTGGCTGCCACGAATATCCAGGCCTTCGCGGTACGCGCAAACGGGCTTTCTGGAACACCTTGCATAG

primer9

GGGCATATGAGCCGTAGCGA

primer10

GCGCCATGGTTTACGGAAAG